

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: March 2, 2002, 20:28:56 ; Search time 2699.18 Seconds
(Without alignments)
2649.239 Million cell updates/sec

Title: US-09-540-235-1
Perfect score: 408
Sequence: 1 acgcgcgtcatgagaatga.....acatcatcatatgttgta 408

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 17159718 seqs, 8763200856 residues

Total number of hits satisfying chosen parameters: 34319436

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database :

Pending_Patents_NA_Main: *
1: /cgn2_6/ptodata/1/pna/US06.COMB.seq: *
2: /cgn2_6/ptodata/1/pna/US06.COMB.seq: *
3: /cgn2_6/ptodata/1/pna/US07.COMB.seq: *
4: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
5: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
6: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
7: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
8: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
9: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
10: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
11: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
12: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
13: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
14: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
15: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
16: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
17: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
18: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
19: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
20: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
21: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
22: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
23: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
24: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
25: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
26: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
27: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
28: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
29: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
30: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
31: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
32: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
33: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
34: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
35: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
36: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
37: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
38: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
39: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
40: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
41: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
42: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *
43: /cgn2_6/ptodata/1/pna/US08.COMB.seq: *

SUMMARIES

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Result No.	Score	Query Match	Length	DB ID	Description
1	408	100.0	408	21	US-09-540-235-1
2	408	100.0	408	45	US-60-128-439-2
3	302.4	74.1	384	45	US-60-128-439-1188
4	302.4	74.1	417	21	US-09-540-235-5042
5	235.2	57.6	298	45	US-60-128-439-1189
6	235.2	57.6	327	21	US-09-540-235-5043
7	133.2	32.6	562	30	US-09-770-152-388
8	133.2	32.6	571	23	US-09-605-701-18273
9	132.8	32.5	384	18	US-09-411-999-36936
10	132.8	32.5	394	23	US-09-605-701-5338
11	132.8	32.5	429	23	US-09-605-699-11618
12	132.8	32.5	471	23	US-09-605-701-4652
13	132.8	32.5	497	23	US-09-606-755-8558
14	132.8	32.5	543	18	US-09-411-999-32416
15	132.8	32.5	555	23	US-09-605-701-17385
16	132.8	32.5	595	58	US-60-253-457-24434
17	132.8	32.5	946	31	US-09-687-716-19329
18	130	31.9	417	27	US-09-654-617-150388
19	130	31.9	417	27	US-09-684-016-150388
20	130	31.9	449	23	US-09-605-701-17346
21	130	31.9	639	22	US-09-565-309A-56964
22	130	31.9	666	58	US-60-253-378-29237
23	130	31.9	724	25	US-09-654-617-127760
24	130	31.9	724	27	US-09-684-016-127760
25	130	31.9	51860	20	US-09-534-859-980
26	129.8	31.8	423	18	US-09-411-999-36980
27	129.8	31.8	385	29	US-09-724-750-32325
28	129.4	31.7	385	29	US-09-724-750-32325
29	129.4	31.7	385	29	US-09-724-750-32325
30	129	31.6	381	22	US-09-605-701-44416
31	129	31.6	639	22	US-09-565-309A-49589
32	126	30.9	668	28	US-09-705-326-2386
33	124.2	30.4	370	27	US-09-654-617-165153
34	124.2	30.4	370	27	US-09-684-016-165153
35	124.2	30.4	432	27	US-09-654-617-165221
36	124.2	30.4	432	27	US-09-684-016-165221
37	124.2	30.4	469	25	US-09-654-617-153103
38	124.2	30.4	469	25	US-09-684-016-153103
39	124.2	30.4	473	27	US-09-654-617-153505
40	124.2	30.4	473	27	US-09-684-016-153505
41	124.2	30.4	475	25	US-09-654-617-161426

42	124.2	30.4	475	27	US-09-684-016-161426	Sequence 161426,
43	124.2	30.4	804	25	US-09-654-617-121273	Sequence 121273,
44	124.2	30.4	804	27	US-09-684-016-121273	Sequence 121273,
45	124.2	30.4	97711	27	US-09-692-412-90	Sequence 90, Appl

ALIGNMENTS

```
RESULT 1
US-09-540-235-1
; Sequence 1, Application US/09540235
; GENERAL INFORMATION:
; APPLICANT: Fisher, Dane K.
; APPLICANT: Laljudi, Raghunath V.
; TITLE OF INVENTION: Nucleic Acid Sequences from Cyanidium caldarium and Uses
; TITLE OF INVENTION: thereof
; FILE REFERENCE: 38-21(15749)B
; CURRENT APPLICATION NUMBER: US/09/540,235
; PRIOR FILING DATE: 2000-04-03
; PRIOR APPLICATION NUMBER: 60/128,439
; PRIOR FILING DATE: 1999-04-06
; NUMBER OF SEQ ID NOS: 5674
; SEQ ID NO 1
; LENGTH: 408
; TYPE: DNA
; ORGANISM: Cyanidium caldarium
; OTHER INFORMATION: Clone ID: LIB190-001-Q1-E1-A12
US-09-540-235-1
```

Query Match 100.0%; Score 408; DB 21; Length 408;

Best Local Similarity 100.0%; Pred. No. 2.1e-113; Mismatches 0; Indels 0; Gaps 0;

Matches 408; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```
QY 1 acgcccgtcgtatgagaattacgagttgcttcgagctcgcgtgttggttcataatttc 60
   |||
Db 1 acgcccgtcgtatgagaattacgagttgcttcgagctcgcgtgttggttcataatttc 60
QY 61 ttgtctgaaagcgtttgggctgttatcagtcgcgatgaagtatccaaagtgtatcgt 120
   |||
Db 61 ttgtctgaaagcgtttgggctgttatcagtcgcgatgaagtatccaaagtgtatcgt 120
QY 121 cgtctagaagaagcagaagaagcattctcaagcacactctctgtaacagaatc 180
   |||
Db 121 cgtctagaagaagcagaagaagcattctcaagcacactctctgtaacagaatc 180
QY 181 tcatgagtgacacctgtgtccaaaggaactacgtacgaataacagcgttcttacct 240
   |||
Db 181 tcatgagtgacacctgtgtccaaaggaactacgtacgaataacagcgttcttacct 240
QY 241 ttccgaaaagaagcagaagaatcatcattgttcgggagcttcaaggttagaagaagaa 300
   |||
Db 241 ttccgaaaagaagcagaagaatcatcattgttcgggagcttcaaggttagaagaagaa 300
QY 301 ttcaacggttatcagaagaatcatcagatcatatcagagagtgaccagaagaagaa 360
   |||
Db 301 ttcaacggttatcagaagaatcatcagatcatatcagagagtgaccagaagaagaa 360
QY 361 cgaatggaatgactgtacgtgttggaatacatcatcatatgttgtta 408
   |||
Db 361 cgaatggaatgactgtacgtgttggaatacatcatcatatgttgtta 408
```

```
RESULT 2
US-60-128-439-2
; Sequence 2, Application US/60128439
; GENERAL INFORMATION:
; APPLICANT: Fisher, Dane K.
; APPLICANT: Laljudi, Raghunath V.
; TITLE OF INVENTION: Nucleic Acid Sequences from Cyanidium caldarium and Uses
; TITLE OF INVENTION: thereof
; FILE REFERENCE: 38-21(15749)A
```

CURRENT APPLICATION NUMBER: US/60/128,439

CURRENT FILING DATE: 1999-04-06

NUMBER OF SEQ ID NOS: 5661

SEQ ID NO 2

LENGTH: 408

TYPE: DNA

ORGANISM: Cyanidium caldarium

FEATURE:

OTHER INFORMATION: Clone ID: LIB190-001-Q1-E1-A12

US-60-128-439-2

Query Match 100.0%; Score 408; DB 45; Length 408;

Best Local Similarity 100.0%; Pred. No. 2.1e-113; Mismatches 0; Indels 0; Gaps 0;

Matches 408; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```
QY 1 acgcccgtcgtatgagaattacgagttgcttcgagctcgcgtgttggttcataatttc 60
   |||
Db 1 acgcccgtcgtatgagaattacgagttgcttcgagctcgcgtgttggttcataatttc 60
QY 61 ttgtctgaaagcgtttgggctgttatcagtcgcgatgaagtatccaaagtgtatcgt 120
   |||
Db 61 ttgtctgaaagcgtttgggctgttatcagtcgcgatgaagtatccaaagtgtatcgt 120
QY 121 cgtctagaagaagcagaagaagcattctcaagcacactctctgtaacagaatc 180
   |||
Db 121 cgtctagaagaagcagaagaagcattctcaagcacactctctgtaacagaatc 180
QY 181 tcatgagtgacacctgtgtccaaaggaactacgtacgaataacagcgttcttacct 240
   |||
Db 181 tcatgagtgacacctgtgtccaaaggaactacgtacgaataacagcgttcttacct 240
QY 241 ttccgaaaagaagcagaagaatcatcattgttcgggagcttcaaggttagaagaagaa 300
   |||
Db 241 ttccgaaaagaagcagaagaatcatcattgttcgggagcttcaaggttagaagaagaa 300
QY 301 ttcaacggttatcagaagaatcatcagatcatatcagagagtgaccagaagaagaa 360
   |||
Db 301 ttcaacggttatcagaagaatcatcagatcatatcagagagtgaccagaagaagaa 360
QY 361 cgaatggaatgactgtacgtgttggaatacatcatcatatgttgtta 408
   |||
Db 361 cgaatggaatgactgtacgtgttggaatacatcatcatatgttgtta 408
```

```
RESULT 3
US-60-128-439-4188
; Sequence 3, Application US/60128439
; GENERAL INFORMATION:
; APPLICANT: Fisher, Dane K.
; APPLICANT: Laljudi, Raghunath V.
; TITLE OF INVENTION: Nucleic Acid Sequences from Cyanidium caldarium and Uses
; TITLE OF INVENTION: thereof
; FILE REFERENCE: 38-21(15749)A
; CURRENT APPLICATION NUMBER: US/60/128,439
; CURRENT FILING DATE: 1999-04-06
; NUMBER OF SEQ ID NOS: 5661
; SEQ ID NO 4188
; LENGTH: 384
; TYPE: DNA
; ORGANISM: Cyanidium caldarium
; FEATURE:
; OTHER INFORMATION: Clone ID: LIB190-054-Q1-E1-D11
US-60-128-439-4188
```

Query Match 74.1%; Score 302.4; DB 45; Length 384;

Best Local Similarity 91.2%; Pred. No. 2.9e-81; Mismatches 31; Indels 0; Gaps 0;

Matches 321; Conservative 0; Mismatches 31; Indels 0; Gaps 0;

```
QY 57 attcttctgaaagcgtttgggctgttatcagtcgcgatgaagtatccaaagtgtta 116
   |||
Db 1 attcttctgaaagcgtttgggctgttatcagtcgcgatgaagtatccaaagtgtta 60
```



```
Db 32 attcgtgtagagcgagcgagagcgagtagtaattgaccagaactatctcaaggttgg 91
Qy 117 tctgtcttaagaagaacagagagagacattctcaagacattctctgtacagca 176
Db 92 tctgtcttaagaagaacagagagagacattctcaagacattctctgtacagca 151
Qy 177 attactatgtagtacccttctcaaggaactatgtagaataacagaggttctt 236
Db 152 attactatgtagtacccttctcaaggaactatgtagaataacagaggttctt 211
Qy 237 cctatcgaagaagaagaagtagtattcgcgcgagcttcaaggttagaaga 296
Db 212 cctatcgaagaagaagaagtagtattcgcgcgagcttcaaggttagaaga 271
Qy 297 aaagttaacagctgttattcgaagaataatcgatatacatcgagagagtagcag 352
Db 272 ccagtcaccaagctgttattcgaagaataatcgatatacatcgagagagtagcag 327
```

```
RESULT 7
US-09-770-152-388
; Sequence 388, Application US/09770152
; GENERAL INFORMATION:
; APPLICANT: Gorlach, Jörn
; APPLICANT: An, Yong-Qiang
; APPLICANT: Hamilton, Carol M.
; APPLICANT: Price, Jennifer L.
; APPLICANT: Raines, Tracy M.
; APPLICANT: Yu, Yang
; APPLICANT: Rameaka, Joshua G.
; APPLICANT: Page, Amy
; APPLICANT: Matthew, Abraham V.
; APPLICANT: Ledford, Brooke L.
; APPLICANT: Woessner, Jeffrey P.
; APPLICANT: Haas, William David
; APPLICANT: Garcia, Carlos A.
; APPLICANT: Krieker, Maja
; APPLICANT: Slader, Ted
; APPLICANT: Davis, Keith R.
; APPLICANT: Allen, Keith
; APPLICANT: Hoffman, Neil
; APPLICANT: Hurban, Patrick
; TITLE OF INVENTION: Expressed Sequences of Arabidopsis
; FILE REFERENCE: 2025US (PRA-014PRV)
; CURRENT APPLICATION NUMBER: US/09/770.152
; CURRENT FILING DATE: 2001-01-26
; PRIOR APPLICATION NUMBER: 60/178,503
; PRIOR FILING DATE: 2000-01-27
; NUMBER OF SEQ ID NOS: 999
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO: 388
; LENGTH: 562
; TYPE: DNA
; ORGANISM: Arabidopsis thaliana
US-09-770-152-388
```

```
Query Match 32.6%; Score 133.2; DB 30; Length 562;
Best Local Similarity 63.4%; Pred. No. 1.1e-29;
Matches 204; Conservative 0; Mismatches 118; Indels 0; Gaps 0;
```

```
Qy 177 tcaatcgcattgaagattccaaagtgtatcgtcttagaagaagaagaggaaga 146
Db 29 tcaatcgcattgaagattccaaagtgtatcgtcttagaagaagaagaggaaga 88
Qy 147 tacttcaagcacacattctgtacagcaataactatgtagcaccctgttcaagaa 206
Db 89 caattcaagctccttcaagcgtgtgagcggttcttcaatgagctgcgcgttcaagaa 148
Qy 207 ctacgtacgaataacagcgttctgttcttcaatcgaaagaagaagaaatcata 266
```

```
Db 149 ctccgttaacaatatacaacgtagatccatgcgattcttgaagaagcagagtagcaat 208
Qy 267 gtccgcgagcttccaaggttagagaagaagaagttacaacggttattgaagaat 326
Db 209 gtccgtgagcgttccaaggttagagaagaagaaggtgtagcaggtgtatccgcaggtg 268
Qy 327 cggatatactcgaagagtagcacaagaagaagaagaatggaatgtagtacctgtgga 386
Db 269 gtgattcaactcgaagagaatcacacaagggagaagaagttacaacgcaacggttcaagctcga 328
Qy 387 ataatcatcattatgtgtta 408
Db 329 gtcaatgtccttgatgtgtaga 350
```

```
RESULT 8
US-09-605-701-18273
; Sequence 18273, Application US/09605701
; GENERAL INFORMATION:
; APPLICANT: Havukkala, Ilkka
; APPLICANT: Shenk, Michael A.
; TITLE OF INVENTION: Polynucleotides, Material Incorporated
; FILE REFERENCE: 11000.104103
; CURRENT APPLICATION NUMBER: US/09/605,701
; CURRENT FILING DATE: 2000-06-21
; NUMBER OF SEQ ID NOS: 25120
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO: 18273
; LENGTH: 371
; TYPE: DNA
; ORGANISM: Pinus radiata
US-09-605-701-18273
```

```
Query Match 32.5%; Score 132.8; DB 23; Length 371;
Best Local Similarity 63.4%; Pred. No. 1.3e-29;
Matches 203; Conservative 0; Mismatches 117; Indels 0; Gaps 0;
```

```
Qy 88 cagtcgcattgaagattccaaagtgtatcgtcttagaagaagaagcagaggaagcat 147
Db 3 catcagaagtagaattacaatccaaagagtcaggttccaggaagaagcagaggaagcag 62
Qy 148 acttcaagcacacattctgtacagcaataactatgtagtgaacctgttccaaggaac 207
Db 63 attcaactgcccatctagtggtgcgtatttgaatgtagtgcacaccttctctgatt 122
Qy 208 taagtacgaataacagcgttcttcttactatctgaagaagaagaagtagaatacatag 267
Db 123 tgagaacaagtagaagtagtcttgcgtatccgattcgaagaagaagtagaagtagcaggtg 182
Qy 268 ttccgcgagcttccaaggttagagaagaagaagttacaacggttattcgaagaatatac 327
Db 183 taagagagacatttaaggttagagaagaagaaggtgttccaaggttattcgaagaatag 242
Qy 328 ggaatacatcgaagagtagcagagaagaagaagaagtagatgtagtacctgttgga 387
Db 243 tcaatacatgttagcagcattactctgtgagaagaatgtagaacaacggttagatgtgtg 302
Qy 388 tacatcatcattatgtgtt 407
Db 303 tccaccatccaaagtggtt 322
```

```
RESULT 9
US-09-411-999-36936
; Sequence 36936, Application US/09411999
; GENERAL INFORMATION:
; APPLICANT: Havukkala, Ilkka
; APPLICANT: Shenk, Michael Andrew
; TITLE OF INVENTION: Polynucleotides Isolated from plants and
; FILE REFERENCE: 10230
```

```

; CURRENT APPLICATION NUMBER: US/09/411,999
; CURRENT FILING DATE: 1999-10-04
; NUMBER OF SEQ ID NOS: 37897
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 36936
; LENGTH: 384
; TYPE: DNA
; ORGANISM: Pinus radiata
US-09-411-999-36936
```

```

Query Match          32.5%; Score 132.8; DB 18; Length 384;
Best Local Similarity 63.4%; Pred. No. 1.3e-29;
Matches 203; Conservative 0; Mismatches 117; Indels 0; Gaps 0;
```

```

QY 88 cagtcgcacatgaagatcttccaaagtgtctgcgtctctagaagaagcagaagaaacat 147
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 44 catcagatgaatgaatcaatccaaagatcagcagttccaggaagaagccgcaagcac 103
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 148 acttcaagcaccatctctctgtacgacgaatactatcatgtagtgcacccctgtccaaaggac 207
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 104 attcaactgccccatcagtgctgcgcgctatttgatgagtgccaccactgtccctctgatt 163
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 208 tacgtacgaataacagcgctgtcttcttaactatcgaaaagaagacgaatcatatag 267
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 164 tgaagaacagatgaatgctgtcgtatccgattagaaagatgatgaaatcagtcg 223
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 268 ttccgagacgttccaggttagaagaagaatgatacaactgttatcgaaagaatc 327
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 224 tccgaggaacatttaagggttagaagaagttgtccaaagttattcgaaagaatgg 283
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 328 ggatacatatcgagagatgacacagaagaacgaatgaaatgactgtactgtggaa 387
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 284 tcatatcgttgagcgacttactcgttgagaaagtcacacactgtgaatgtgtg 343
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 388 tacatccatctaatgttgt 407
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 344 tccaccatccaaagtcgt 363
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
```

```

RESULT 10
US-09-605-701-5338
; Sequence 5338, Application US/09605701
; GENERAL INFORMATION:
; APPLICANT: Havukkala, Ilkka
; APPLICANT: Shenk, Michael A.
; TITLE OF INVENTION: Polynucleotides, Materials Incorporated
; FILE REFERENCE: 11000.104103
; CURRENT APPLICATION NUMBER: US/09/605,701
; CURRENT FILING DATE: 2000-06-21
; NUMBER OF SEQ ID NOS: 25120
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 5338
; LENGTH: 394
; TYPE: DNA
; ORGANISM: Pinus radiata
US-09-605-701-5338
```

```

Query Match          32.5%; Score 132.8; DB 23; Length 394;
Best Local Similarity 63.4%; Pred. No. 1.3e-29;
Matches 203; Conservative 0; Mismatches 117; Indels 0; Gaps 0;
```

```

QY 88 cagtcgcacatgaagatcttccaaagtgtctgcgtctctagaagaagcagaagaaacat 147
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 53 catcagatgaatgaatcaatccaaagatcagcagttccaggaagaagccgcaagcac 112
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 148 acttcaagcaccatctctgtacgacgaatactatcatgtagtgcacccctgtccaaaggac 207
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 113 attcaactgccccatcagtgctgcgcgctatttgatgagtgccaccactgtccctctgatt 172
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 208 tacgtacgaataacagcgctgtcttcttaactatcgaaaagaagacgaatcatatg 267
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
```

```

DB 173 tgaagaacagatgaatcgttctgtccatccgattagaagaagatgaagatcagtg 232
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 268 ttccgagacgttccaaaggtagaagaagaatgatacaactgttatcgaaagaatc 327
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 233 tacgaggaacatttaagggttagaagaagaagttgtccaaagttatcgaaagaatgg 292
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 328 ggatacatatcgagagatgacacagaagaacgaatgaaatgactgtactgtggaa 387
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 293 tcatatcgttgagcgacttactcgttgagaaagtcacacactgtgaatgtgtg 352
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 388 tacatccatctaatgttgt 407
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 353 tccaccatccaaagtcgt 372
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
```

```

RESULT 11
US-09-605-699-11618
; Sequence 11618, Application US/09605699
; GENERAL INFORMATION:
; APPLICANT: Havukkala, Ilkka
; APPLICANT: Shenk, Michael A.
; TITLE OF INVENTION: Polynucleotides, Materials Incorporated
; FILE REFERENCE: 11000.104102
; CURRENT APPLICATION NUMBER: US/09/605,699
; CURRENT FILING DATE: 2000-06-21
; NUMBER OF SEQ ID NOS: 25120
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 11618
; LENGTH: 429
; TYPE: DNA
; ORGANISM: Pinus radiata
US-09-605-699-11618
```

```

Query Match          32.5%; Score 132.8; DB 23; Length 429;
Best Local Similarity 63.4%; Pred. No. 1.3e-29;
Matches 203; Conservative 0; Mismatches 117; Indels 0; Gaps 0;
```

```

QY 88 cagtcgcacatgaagatcttccaaagtgtctgcgtctctagaagaagcagaagaaacat 147
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 53 catcagatgaatgaatcaatccaaagatcagcagttccaggaagaagccgcaagcac 112
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 148 acttcaagcaccatctctgtacgacgaatactatcatgtagtgcacccctgtccaaaggac 207
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 113 attcaactgccccatcagtgctgcgcgctatttgatgagtgccaccactgtccctctgatt 172
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 208 tacgtacgaataacagcgctgtcttcttaactatcgaaaagaagacgaatcatatg 267
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 173 tgaagaacagatgaatcgttctgtccatccgattagaagaagatgaagttcagtg 232
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 268 ttccgagacgttccaaaggtagaagaagaatgatacaactgttatcgaaagaatc 327
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 233 tacgaggaacatttaagggtagaagaaggtgttccaaagttatcgaaagaatgg 292
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 328 ggatacatatcgagagatgacacagaagaacgaatgaaatgactgtactgtggaa 387
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 293 tcatatcgttgagcgacttactcgttgagaaagtcacacactgtgaatgtgtg 352
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
QY 388 tacatccatctaatgttgt 407
   ||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 353 tccaccatccaaagtcgt 372
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
```

```

RESULT 12
US-09-605-701-4652
; Sequence 4652, Application US/09605701
; GENERAL INFORMATION:
; APPLICANT: Havukkala, Ilkka
; APPLICANT: Shenk, Michael A.
; TITLE OF INVENTION: Polynucleotides, Materials Incorporated
; FILE REFERENCE: 11000.104103
; CURRENT APPLICATION NUMBER: US/09/605,701
; CURRENT FILING DATE: 2000-06-21
; NUMBER OF SEQ ID NOS: 25120
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 4652
; LENGTH: 394
; TYPE: DNA
; ORGANISM: Pinus radiata
US-09-605-701-4652
```

```
FILE REFERENCE: 11000.1041U3
; CURRENT APPLICATION NUMBER: US/09/605.701
; NUMBER FILING DATE: 2000-06-21
; NUMBER OF SEQ ID NOS: 25120
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 4652
; LENGTH: 471
; TYPE: DNA
; ORGANISM: Pinus radiata
US-09-605-701-4652
```

```
Query Match          32.5%; Score 132.8; DB 23; Length 471;
Best Local Similarity 63.4%; Pred. No. 1.4e-29;
Matches 203; Conservative 0; Mismatches 117; Indels 0; Gaps 0;
```

```
QY 88 cagtcgcataagatcccaagtgatcgtctctagaagaagcagaagcat 147
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 78 catcaagatgaagtaacatccaagagtcagcagttccaggaagaagcgccgaagcac 137
QY 148 acttcaagcaccatctctctgaagcaatactatgaagcaccctgtccaaagcac 207
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 138 attcaacgcccactagtgctgcgtatttgatgaagtcaccactgtctctgatt 197
QY 208 tacgtaagaatacagcgctgtctcttaccattcgaagaagcaagcaatcatcag 267
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 198 tgagaacaagaacaaatgtctctgcattcagatagaagaagatgaagttcagtg 257
QY 268 ttccgagcgtcttcaagggtagaagaagaattacaacgtgtatcgaaagaatc 327
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 258 tacgaggaacatttaagggtagaagaagttgtccaaagttatccgaagaatg 317
QY 328 gatacatatcgagagtgagcagaagaagaagcaatgaatgaatgaatgaatg 387
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 318 tcatcatatgttgagcagctactcgtgagaagaagtcacacgtgaaatgtgtg 377
QY 388 tacatccatcaatgtgtgt 407
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 378 tccaccatcccaagtgctgt 397
```

```
RESULT 13
US-09-606-755-8558
; Sequence 8558, Application US/09606755
; GENERAL INFORMATION:
; APPLICANT: Havukkala, Ilkka
; APPLICANT: Shenk, Michael A.
; TITLE OF INVENTION: Polynucleotides, Materials Incorporating
; FILE REFERENCE: 11000.1041U6
; CURRENT APPLICATION NUMBER: US/09/606.755
; CURRENT FILING DATE: 2000-06-22
; NUMBER OF SEQ ID NOS: 17747
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8558
; LENGTH: 497
; TYPE: DNA
; ORGANISM: Pinus radiata
US-09-606-755-8558
```

```
Query Match          32.5%; Score 132.8; DB 23; Length 497;
Best Local Similarity 63.4%; Pred. No. 1.4e-29;
Matches 203; Conservative 0; Mismatches 117; Indels 0; Gaps 0;
```

```
QY 88 cagtcgcataagatcccaagtgatcgtctctagaagaagcagaagcat 147
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 26 catcaagatgaagtaacatccaagagtcagcagttccaggaagaagcgccgaagcac 85
QY 148 acttcaagcaccatctctctgaagcaatactatgaagcaccctgtccaaagcac 207
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 86 attcaacgcccactagtgctgcgtatttgatgaagtcaccactgtctctgatt 145
```

```
QY 208 tacgtcaaatatacagcgtctcttcttactatccgaagaagaagcaagtaatcatg 267
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 146 tggaaacaagaatgaatgtctgtcttccgttactgaagaagatgaatgaatgagtg 205
QY 268 ttccgagagcttcaagggtagaagaagaattacaacggtttatcgaaagaatc 327
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 206 tacgaggaacatttaagggtagaagaagttgtccaaagttatcgaaagaatg 265
QY 328 gatacatatcgagagtgagcagaagaagaagcaatgaatgaatgaatgaatg 387
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 266 tcatcatatgttgagcagctactcgtgagaagaatgcacacgtgaaatgtgtg 325
QY 388 tacatccatcaatgtgtgt 407
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 326 tccaccatcccaagtgctgt 345
```

```
RESULT 14
US-09-411-999-32416
; Sequence 32416, Application US/09411999
; GENERAL INFORMATION:
; APPLICANT: Havukkala, Ilkka
; APPLICANT: Shenk, Michael Andrew
; TITLE OF INVENTION: Polynucleotides isolated from plants and
; FILE REFERENCE: 1023U
; CURRENT APPLICATION NUMBER: US/09/411.999
; CURRENT FILING DATE: 1999-10-04
; NUMBER OF SEQ ID NOS: 37897
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 32416
; LENGTH: 543
; TYPE: DNA
; ORGANISM: Pinus radiata
US-09-411-999-32416
```

```
Query Match          32.5%; Score 132.8; DB 18; Length 543;
Best Local Similarity 63.4%; Pred. No. 1.4e-29;
Matches 203; Conservative 0; Mismatches 117; Indels 0; Gaps 0;
```

```
QY 88 cagtcgcataagatcccaagtgatcgtctctagaagaagcagaagcat 147
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 71 catcaagatgaagtaacatccaagagtcagcagttccaggaagaagcgccgaagcac 130
QY 148 acttcaagcaccatctctctgaagcaatactatgaagcaccctgtccaaagcac 207
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 131 attcaacgcccactagtgctgcgtatttgatgaagtcaccactgtctctgatt 190
QY 208 tacgtcaaatatacagcgtctcttactatccgaagaagaagcaatgaatcatg 267
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 191 tgagaacaagaatgaatgtctgtcgtatccgaatagaagaagatgaatgaatg 250
QY 268 ttccgagagcttcaagggtagaagaagaattacaacgtgttatcgaaagaatc 327
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 251 tacgaggaacatttaagggtagaagaagttgtccaaagttatccgaagaatg 310
QY 328 gatacatatcgagagtgagcagaagaagaagcaatgaatgaatgaatgaatg 387
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 311 tcatcatatgttgagcagctactcgtgagaagaatgcacacgtgaaatgtgtg 370
QY 388 tacatccatcaatgtgtgt 407
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 371 tccaccatcccaagtgctgt 390
```

```
RESULT 15
US-09-605-701-17385
; Sequence 17385, Application US/09605701
; GENERAL INFORMATION:
; APPLICANT: Havukkala, Ilkka
; APPLICANT: Shenk, Michael A.
; TITLE OF INVENTION: Polynucleotides, Material Incorporated
```

```
; TITLE OF INVENTION: Them, and Methods for Using Them
; FILE REFERENCE: 11000.1041u3
; CURRENT APPLICATION NUMBER: US/09/605,701
; CURRENT FILING DATE: 2000-06-21
; NUMBER OF SEQ ID NOS: 25120
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 17385
; LENGTH: 555
; TYPE: DNA
; ORGANISM: Pinus radiata
US-09-605-701-17385
```

```
Query Match          32.5%; Score 132.8; DB 23; Length 555;
Best Local Similarity 63.4%; Pred No.1.5e-29;
Matches 203; Conservative 0; Mismatches 117; Indels 0; Gaps 0;

QY 88 cagtcgcacatgaagatcccaagtgtatcgtcgtctagaagaagcagagaagcat 147
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 69 catcagatgaagtacatccaagtcagcagttccagagaagagccgaagcac 128

QY 148 acttcaagcacatctctgtacagcaatactcatgtacgacctgtccaaggaa 207
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 123 attcacgtcccatctagtgctggtgtatcttgatgtgtcaccactgtcctctgalt 188

QY 208 tacgtacgaatacagcgcttctcttacctatcgaaagaagacgaatcatcatg 267
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 189 tgagaacaagtaacaatgtctgtcgtatccgattcagaagaagatgaatgcaagtgg 248

QY 268 ttcggtgagcttcaagggtagaagaagaagtacaaagtgtatcgaaagaataatc 327
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 249 tacgaagaacatttaagggtagaagaaggtgtcccaagttatcgaagaagaatggg 308

QY 328 ggatacatatcgagagagtgaccagagaagaagcgaatggaatgactgtgggaa 387
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 309 tcatacatgttagcgcaattactcgttgagaagaatcaatgycacaactgtgaatgtgtg 368

QY 388 tacatcatcatatgttgt 407
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 369 tccaccatccaaggtcgt 388
```

Search completed: March 2, 2002, 23:02:24
Job time: 9208 sec

